

## SEQUENCE LISTING

<110> AZPIROZ, Ricardo  
CHOE, Sunghwa  
FELDMANN, Kenneth

<120> DWF4 POLYNUCLEOTIDES, POLYPEPTIDES AND USES THEREOF

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 Thr Leu Phe Val Leu Asp Ser Trp Gln Gln Asn Ser Ile Phe Ser Ala  
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Ala	Met	Thr	Asp	Val	Glu	Ile	Lys	Gly	Tyr	Lys	Ile	Pro	Lys	Gly	Trp		
	355					360						365					
Lys	Val	Phe	Ser	Ser	Phe	Arg	Ala	Val	His	Leu	Asp	Pro	Asn	His	Phe		
	370					375					380						
Lys	Asp	Ala	Arg	Thr	Phe	Asn	Pro	Trp	Arg	Trp	Gln	Ser	Asn	Ser	Val		
385				390						395					400		
Thr	Thr	Gly	Pro	Ser	Asn	Val	Phe	Thr	Pro	Phe	Gly	Gly	Gly	Pro	Arg		
			405					410						415			
Leu	Cys	Pro	Gly	Tyr	Glu	Leu	Ala	Arg	Val	Ala	Leu	Ser	Val	Phe	Leu		
		420						425					430				
His	Arg	Leu	Val	Thr	Gly	Phe	Ser	Trp	Val	Pro	Ala	Glu	Gln	Asp	Lys		
	435						440				445						
Leu	Val	Phe	Phe	Pro	Thr	Thr	Arg	Thr	Gln	Lys	Arg	Tyr	Pro	Ile	Phe		
	450					455					460						
Val	Lys	Arg	Arg	Asp	Phe	Ala	Thr										
465					470												

&lt;210&gt; 20

&lt;211&gt; 464

&lt;212&gt; PRT

&lt;213&gt; Lycopersicon esculentum

&lt;400&gt; 20

Met	Ala	Phe	Phe	Leu	Ile	Phe	Leu	Ser	Ser	Phe	Phe	Gly	Leu	Cys	Ile		
1				5				10					15				
Phe	Cys	Thr	Ala	Leu	Leu	Arg	Trp	Asn	Gln	Val	Lys	Tyr	Asn	Gln	Lys		
		20					25					30					
Asn	Leu	Pro	Pro	Gly	Thr	Met	Gly	Trp	Pro	Leu	Phe	Gly	Glu	Thr	Thr		
	35					40						45					
Glu	Phe	Leu	Lys	Leu	Gly	Pro	Ser	Phe	Met	Lys	Asn	Gln	Arg	Ala	Arg		
	50					55				60							
Tyr	Gly	Ser	Phe	Phe	Lys	Ser	His	Ile	Leu	Gly	Cys	Pro	Thr	Ile	Val		
65					70				75					80			
Ser	Met	Asp	Ser	Glu	Leu	Asn	Arg	Tyr	Ile	Leu	Val	Asn	Glu	Ala	Lys		
			85					90				95					
Gly	Leu	Val	Pro	Gly	Tyr	Pro	Gln	Ser	Met	Ile	Asp	Ile	Leu	Gly	Lys		

			100				105					110				
Cys	Asn	Ile	Ala	Ala	Val	Asn	Gly	Ser	Ala	His	Lys	Tyr	Met	Arg	Gly	
			115				120					125				
Ala	Leu	Leu	Ser	Leu	Ile	Ser	Pro	Thr	Met	Ile	Arg	Asp	Gln	Leu	Leu	
			130				135					140				
Pro	Lys	Ile	Asp	Glu	Phe	Met	Arg	Ser	His	Leu	Thr	Asn	Trp	Asp	Asn	
145				150				155					160			
Lys	Val	Ile	Asp	Ile	Gln	Glu	Lys	Thr	Asn	Lys	Met	Ala	Phe	Leu	Ser	
			165				170					175				
Ser	Leu	Lys	Gln	Ile	Ala	Gly	Ile	Glu	Ser	Thr	Ser	Leu	Ala	Gln	Glu	
			180				185					190				
Phe	Met	Ser	Glu	Phe	Phe	Asn	Leu	Val	Leu	Gly	Thr	Leu	Ser	Leu	Pro	
			195				200					205				
Ile	Asn	Leu	Pro	Asn	Thr	Asn	Tyr	His	Arg	Gly	Phe	Gln	Ala	Arg	Lys	
			210				215					220				
Ile	Ile	Val	Asn	Leu	Leu	Arg	Thr	Leu	Ile	Glu	Glu	Arg	Arg	Ala	Ser	
225				230				235					240			
Lys	Glu	Ile	Gln	His	Asp	Met	Leu	Gly	Tyr	Leu	Met	Asn	Glu	Glu	Ala	
			245				250					255				
Thr	Arg	Phe	Lys	Leu	Thr	Asp	Asp	Glu	Met	Ile	Asp	Leu	Ile	Ile	Thr	
			260				265					270				
Ile	Leu	Tyr	Ser	Gly	Tyr	Glu	Thr	Val	Ser	Thr	Thr	Ser	Met	Met	Ala	
			275				280					285				
Val	Lys	Tyr	Leu	His	Asp	His	Pro	Lys	Val	Leu	Glu	Glu	Leu	Arg	Lys	
			290				295					300				
Glu	His	Met	Ala	Ile	Arg	Glu	Lys	Lys	Lys	Pro	Glu	Asp	Pro	Ile	Asp	
305				310				315					320			
Tyr	Asn	Asp	Tyr	Arg	Ser	Met	Arg	Phe	Thr	Arg	Ala	Val	Ile	Leu	Glu	
			325				330					335				
Thr	Ser	Arg	Leu	Ala	Thr	Ile	Val	Asn	Gly	Val	Leu	Arg	Lys	Thr	Thr	
			340				345					350				
Gln	Asp	Met	Glu	Ile	Asn	Gly	Tyr	Ile	Ile	Pro	Lys	Gly	Trp	Arg	Ile	
			355				360					365				
Tyr	Val	Tyr	Thr	Arg	Glu	Leu	Asn	Tyr	Asp	Pro	Arg	Leu	Tyr	Pro	Asp	
			370				375					380				
Pro	Tyr	Ser	Phe	Asn	Pro	Trp	Arg	Trp	Met	Asp	Lys	Ser	Leu	Glu	His	
385				390				395					400			
Gln	Asn	Ser	Phe	Leu	Val	Phe	Gly	Gly	Gly	Thr	Arg	Gln	Cys	Pro	Gly	
			405				410					415				
Lys	Glu	Leu	Gly	Val	Ala	Glu	Ile	Ser	Thr	Phe	Leu	His	Tyr	Phe	Val	
			420				425					430				
Thr	Lys	Tyr	Arg	Trp	Glu	Glu	Ile	Gly	Gly	Asp	Lys	Leu	Met	Lys	Phe	
			435				440					445				
Pro	Arg	Val	Glu	Ala	Pro	Asn	Gly	Leu	Arg	Ile	Arg	Val	Ser	Ala	His	
			450				455					460				

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<210> 21
<211> 444
<212> PRT
<213> Synechocystis sp.
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<400> 21															
Met	Ile	Thr	Ser	Pro	Thr	Asn	Leu	Asn	Ser	Leu	Pro	Ile	Pro	Pro	Gly
1				5					10					15	
Asp	Phe	Gly	Leu	Pro	Trp	Leu	Gly	Glu	Thr	Leu	Asn	Phe	Leu	Asn	Asp
			20					25					30		
Gly	Asp	Phe	Gly	Lys	Lys	Arg	Gln	Gln	Gln	Phe	Gly	Pro	Ile	Phe	Lys

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<210> 22
<211> 519
<212> PRT
<213> Zea mays
```

<400> 22

Met	Leu	Gly	Val	Gly	Met	Ala	Ala	Ala	Val	Leu	Leu	Gly	Ala	Val	Ala
1				5					10					15	
Leu	Leu	Leu	Ala	Asp	Ala	Ala	Ala	Arg	Arg	Ala	His	Trp	Trp	Tyr	Arg
			20					25					30		
Glu	Ala	Ala	Glu	Ala	Val	Leu	Val	Gly	Ala	Val	Ala	Leu	Val	Val	Val
			35					40					45		
Asp	Ala	Ala	Ala	Arg	Arg	Ala	His	Gly	Trp	Tyr	Arg	Glu	Ala	Ala	Leu
			50				55				60				
Gly	Ala	Ala	Arg	Arg	Ala	Arg	Leu	Pro	Pro	Gly	Glu	Met	Gly	Trp	Pro
65					70					75					80
Leu	Val	Gly	Gly	Met	Trp	Ala	Phe	Leu	Arg	Ala	Phe	Lys	Ser	Gly	Lys
				85					90					95	
Pro	Asp	Ala	Phe	Ile	Ala	Ser	Phe	Val	Arg	Arg	Phe	Gly	Arg	Thr	Gly
			100					105					110		
Val	Tyr	Arg	Ser	Phe	Met	Phe	Ser	Ser	Pro	Thr	Val	Leu	Val	Thr	Thr
			115					120					125		
Ala	Glu	Gly	Cys	Lys	Gln	Val	Leu	Met	Asp	Asp	Asp	Ala	Phe	Val	Thr
			130				135					140			
Gly	Trp	Pro	Lys	Ala	Thr	Val	Ala	Leu	Val	Gly	Pro	Arg	Ser	Phe	Val
145					150					155					160
Ala	Met	Pro	Tyr	Asp	Glu	His	Arg	Arg	Ile	Arg	Lys	Leu	Thr	Ala	Ala
				165					170					175	
Pro	Ile	Asn	Gly	Phe	Asp	Ala	Leu	Thr	Gly	Tyr	Leu	Pro	Phe	Ile	Asp
			180					185					190		
Arg	Thr	Val	Thr	Ser	Ser	Leu	Arg	Ala	Trp	Ala	Asp	His	Gly	Gly	Ser
		195					200					205			
Val	Glu	Phe	Leu	Thr	Glu	Leu	Arg	Arg	Met	Thr	Phe	Lys	Ile	Ile	Val
		210					215					220			
Gln	Ile	Phe	Leu	Gly	Gly	Ala	Asp	Gln	Ala	Thr	Thr	Arg	Ala	Leu	Glu
225					230					235					240
Arg	Ser	Tyr	Thr	Glu	Leu	Asn	Tyr	Gly	Met	Arg	Ala	Met	Ala	Ile	Asn
				245					250					255	
Leu	Pro	Gly	Phe	Ala	Tyr	Arg	Gly	Ala	Leu	Arg	Ala	Arg	Arg	Arg	Leu
			260					265					270		
Val	Ala	Val	Leu	Gln	Gly	Val	Leu	Asp	Glu	Arg	Arg	Ala	Ala	Arg	Ala
			275					280				285			
Lys	Gly	Val	Ser	Gly	Gly	Gly	Val	Asp	Met	Met	Asp	Arg	Leu	Ile	Glu
			290			295					300				
Ala	Gln	Asp	Glu	Arg	Gly	Arg	His	Leu	Asp	Asp	Asp	Glu	Ile	Ile	Asp
305					310					315					320
Val	Leu	Val	Met	Tyr	Leu	Asn	Ala	Gly	His	Glu	Ser	Ser	Gly	His	Ile
				325					330					335	
Thr	Met	Trp	Ala	Thr	Val	Phe	Leu	Gln	Glu	Asn	Pro	Asp	Met	Phe	Ala
			340					345					350		
Arg	Ala	Lys	Ala	Glu	Gln	Glu	Ala	Ile	Met	Arg	Ser	Ile	Pro	Ser	Ser
			355				360					365			
Gln	Arg	Gly	Leu	Thr	Leu	Arg	Asp	Phe	Arg	Lys	Met	Glu	Tyr	Leu	Ser
			370				375				380				
Gln	Val	Ile	Asp	Glu	Thr	Leu	Arg	Leu	Val	Asn	Ile	Ser	Phe	Val	Ser
385					390					395					400
Phe	Arg	Gln	Ala	Thr	Arg	Asp	Val	Phe	Val	Asn	Gly	Tyr	Leu	Ile	Pro
				405					410					415	
Lys	Gly	Trp	Lys	Val	Gln	Leu	Trp	Tyr	Arg	Ser	Val	His	Met	Asp	Pro
			420					425					430		
Gln	Val	Tyr	Pro	Asp	Pro	Thr	Lys	Phe	Asp	Pro	Ser	Arg	Trp	Glu	Gly
			435				440						445		

His Ser Pro Arg Ala Gly Thr Phe Leu Ala Phe Gly Leu Gly Ala Arg  
 450 455 460  
 Leu Cys Pro Gly Asn Asp Leu Ala Lys Leu Glu Ile Ser Val Phe Leu  
 465 470 475 480  
 His His Phe Leu Leu Gly Tyr Lys Leu Ala Arg Thr Asn Pro Arg Cys  
 485 490 495  
 Arg Val Arg Tyr Leu Pro His Pro Arg Pro Val Asp Asn Cys Leu Ala  
 500 505 510  
 Lys Ile Thr Arg Val Gly Ser  
 515

<210> 23

<211> 492

<212> PRT

<213> Danio rerio

<400> 23

Met Gly Leu Tyr Thr Leu Met Val Thr Phe Leu Cys Thr Ile Val Leu  
 1 5 10 15  
 Pro Val Leu Leu Phe Leu Ala Ala Val Lys Leu Trp Glu Met Leu Met  
 20 25 30  
 Ile Arg Arg Val Asp Pro Asn Cys Arg Ser Pro Leu Pro Pro Gly Thr  
 35 40 45  
 Met Gly Leu Pro Phe Ile Gly Glu Thr Leu Gln Leu Ile Leu Gln Arg  
 50 55 60  
 Arg Lys Phe Leu Arg Met Lys Arg Gln Lys Tyr Gly Cys Ile Tyr Lys  
 65 70 75 80  
 Thr His Leu Phe Gly Asn Pro Thr Val Arg Val Met Gly Ala Asp Asn  
 85 90 95  
 Val Arg Gln Ile Leu Leu Gly Glu His Lys Leu Val Ser Val Gln Trp  
 100 105 110  
 Pro Ala Ser Val Arg Thr Ile Leu Gly Ser Asp Thr Leu Ser Asn Val  
 115 120 125  
 His Gly Val Gln His Lys Asn Lys Lys Lys Ala Ile Met Arg Ala Phe  
 130 135 140  
 Ser Arg Asp Ala Leu Glu His Tyr Ile Pro Val Ile Gln Gln Glu Val  
 145 150 155 160  
 Lys Ser Ala Ile Gln Glu Trp Leu Gln Lys Asp Ser Cys Val Leu Val  
 165 170 175  
 Tyr Pro Glu Met Lys Lys Leu Met Phe Arg Ile Ala Met Arg Ile Leu  
 180 185 190  
 Leu Gly Phe Glu Pro Glu Gln Ile Lys Thr Asp Glu Gln Glu Leu Val  
 195 200 205  
 Glu Ala Phe Glu Glu Met Ile Lys Asn Leu Phe Ser Leu Pro Ile Asp  
 210 215 220  
 Val Pro Phe Ser Gly Leu Tyr Arg Gly Leu Arg Ala Arg Asn Phe Ile  
 225 230 235 240  
 His Ser Lys Ile Glu Glu Asn Ile Arg Lys Lys Ile Gln Asp Asp Asp  
 245 250 255  
 Asn Glu Asn Glu Gln Lys Tyr Lys Asp Ala Leu Gln Leu Leu Ile Glu  
 260 265 270  
 Asn Ser Arg Arg Ser Asp Glu Pro Phe Ser Leu Gln Ala Met Lys Glu  
 275 280 285  
 Ala Ala Thr Glu Leu Leu Phe Gly Gly His Glu Thr Thr Ala Ser Thr  
 290 295 300  
 Ala Thr Ser Leu Val Met Phe Leu Gly Leu Asn Thr Glu Val Val Gln  
 305 310 315 320

```

Lys Val Arg Glu Glu Val Gln Glu Lys Val Glu Met Gly Met Tyr Thr
          325          330          335
Pro Gly Lys Gly Leu Ser Met Glu Leu Leu Asp Gln Leu Lys Tyr Thr
          340          345          350
Gly Cys Val Ile Lys Glu Thr Leu Arg Ile Asn Pro Pro Val Pro Gly
          355          360          365
Gly Phe Arg Val Ala Leu Lys Thr Phe Glu Leu Asn Gly Tyr Gln Ile
          370          375          380
Pro Lys Gly Trp Asn Val Ile Tyr Ser Ile Cys Asp Thr His Asp Val
385          390          395          400
Ala Asp Val Phe Pro Asn Lys Glu Glu Phe Gln Pro Glu Arg Phe Met
          405          410          415
Ser Lys Gly Leu Glu Asp Gly Ser Arg Phe Asn Tyr Ile Pro Phe Gly
          420          425          430
Gly Gly Ser Arg Met Cys Val Gly Lys Glu Phe Ala Lys Val Leu Leu
          435          440          445
Lys Ile Phe Leu Val Glu Leu Thr Gln His Cys Asn Trp Ile Leu Ser
          450          455          460
Asn Gly Pro Pro Thr Met Lys Thr Gly Pro Thr Ile Tyr Pro Val Asp
465          470          475          480
Asn Leu Pro Thr Lys Phe Thr Ser Tyr Val Arg Asn
          485          490

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&lt;210&gt; 24

&lt;211&gt; 504

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 24

```

Met Ala Leu Ile Pro Asp Leu Ala Met Glu Thr Trp Leu Leu Leu Ala
1          5          10          15
Val Ser Leu Val Leu Leu Tyr Leu Tyr Gly Thr His Ser His Gly Leu
          20          25          30
Phe Lys Lys Leu Gly Ile Pro Gly Pro Thr Pro Leu Pro Phe Leu Gly
          35          40          45
Asn Ile Leu Ser Tyr His Lys Gly Phe Cys Met Phe Asp Met Glu Cys
          50          55          60
His Lys Lys Tyr Gly Lys Val Trp Gly Phe Tyr Asp Gly Gln Gln Pro
65          70          75          80
Val Leu Ala Ile Thr Asp Pro Asp Met Ile Lys Leu Val Leu Val Lys
          85          90          95
Glu Cys Tyr Ser Val Phe Thr Asn Arg Glu Pro Phe Gly Pro Val Gly
          100          105          110
Phe Met Lys Ser Ala Ile Ser Ile Ala Glu Asp Glu Glu Trp Lys Arg
          115          120          125
Leu Arg Ser Leu Leu Ser Pro Thr Phe Thr Ser Gly Lys Leu Lys Glu
          130          135          140
Met Val Pro Ile Ile Ala Gln Tyr Gly Asp Val Leu Val Arg Asn Leu
145          150          155          160
Arg Arg Glu Arg Glu Thr Gly Lys Pro Val Thr Leu Lys Asp Val Phe
          165          170          175
Gly Ala Tyr Ser Met Asp Val Ile Thr Ser Ser Ser Phe Gly Val Asn
          180          185          190
Val Asp Ser Leu Asn Asn Pro Gln Asp Pro Leu Val Glu Asn Thr Lys
          195          200          205
Lys Leu Leu Arg Phe Asp Phe Leu Asp Pro Phe Phe Leu Ser Ile Thr
210          215          220

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Val Phe Pro Phe Leu Ile Pro Ile Leu Glu Val Leu Asn Ile Cys Val
225                230                235                240
Phe Pro Arg Glu Val Thr Asn Phe Leu Arg Lys Ala Val Lys Arg Met
                245                250                255
Lys Glu Ser Arg Leu Glu Asp Thr Gln Lys His Arg Val Asp Phe Leu
                260                265                270
Gln Leu Met Ile Asp Ser His Lys Asn Ser Lys Glu Thr Glu Ser His
                275                280                285
Lys Ala Leu Ser Asp Leu Glu Leu Val Ala Gln Ser Ile Ile Phe Ile
                290                295                300
Phe Ala Gly Tyr Glu Thr Thr Ser Ser Val Leu Ser Phe Ile Met Tyr
305                310                315                320
Glu Leu Ala Thr His Pro Asp Val Gln Gln Lys Leu Gln Glu Glu Ile
                325                330                335
Asp Ala Val Leu Pro Asn Lys Ala Pro Pro Thr Tyr Asp Thr Val Leu
                340                345                350
Gln Met Glu Tyr Leu Asp Met Val Val Asn Glu Thr Leu Arg Leu Phe
                355                360                365
Pro Ile Ala Met Arg Leu Glu Arg Val Cys Lys Lys Asp Val Glu Ile
                370                375                380
Asn Gly Met Phe Ile Pro Lys Gly Trp Val Val Met Ile Pro Ser Tyr
385                390                395                400
Ala Leu His Arg Asp Pro Lys Tyr Trp Thr Glu Pro Glu Lys Phe Leu
                405                410                415
Pro Glu Arg Phe Ser Lys Lys Asn Lys Asp Asn Ile Asp Pro Tyr Ile
                420                425                430
Tyr Thr Pro Phe Gly Ser Gly Pro Arg Asn Cys Ile Gly Met Arg Phe
                435                440                445
Ala Leu Met Asn Met Lys Leu Ala Leu Ile Arg Val Leu Gln Asn Phe
                450                455                460
Ser Phe Lys Pro Cys Lys Glu Thr Gln Ile Pro Leu Lys Leu Ser Leu
465                470                475                480
Gly Gly Leu Leu Gln Pro Glu Lys Pro Val Val Leu Lys Val Glu Ser
                485                490                495
Arg Asp Gly Thr Val Ser Gly Ala
                500

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<210> 25

<211> 575

<212> PRT

<213> Artificial Sequence

<220>

<223> Consensus sequence

<221> VARIANT

<222> (1)...(575)

<223> Xaa = Any Amino Acid or No Amino Acid

<400> 25

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1                5                10                15
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                20                25                30
Xaa Xaa Xaa Xaa Xaa Xaa Leu Leu Ser Xaa Xaa Ala Leu Xaa Val Xaa
                35                40                45
Leu Xaa Leu Ala Ala Arg Arg Xaa Xaa Xaa Arg Tyr Xaa Xaa Xaa Xaa

```

	50				55				60						
Xaa 65	Xaa	Xaa	Xaa	Arg	Arg	Lys	Xaa	Leu	Pro	Pro	Gly	Thr	Met	Gly	Leu
Pro	Xaa	Leu	Gly	Glu	Thr	Leu	Gln	Phe	Leu	Lys	Xaa	Xaa	Xaa	Xaa	Xaa
				85					90					95	
Xaa	Pro	Gly	Asp	Phe	Xaa	Lys	Glu	Arg	Val	Xaa	Xaa	Tyr	Gly	Xaa	Xaa
			100					105					110		
Xaa	Xaa	Ile	Tyr	Lys	His	Leu	Phe	Gly	Glu	Pro	Thr	Ile	Xaa	Ser	Xaa
		115					120					125			
Asp	Ala	Glu	Leu	Asn	Arg	Phe	Xaa	Leu	Xaa	Asn	Glu	Gly	Xaa	Lys	Leu
	130					135					140				
Phe	Xaa	Cys	Xaa	Xaa	Pro	Ala	Ser	Xaa	Xaa	Gly	Xaa	Leu	Gly	Lys	Xaa
145					150					155					160
Ser	Leu	Xaa	Ala	Xaa	Xaa	Gly	Xaa	Glu	His	Lys	Arg	Met	Arg	Xaa	Leu
				165					170					175	
Leu	Xaa	Ser	Xaa	Phe	Ser	Xaa	Xaa	Xaa	Xaa	Leu	Asp	His	Xaa	Leu	Pro
			180					185					190		
Xaa	Ile	Asp	Arg	Xaa	Val	Arg	Ser	Xaa	Leu	Xaa	Xaa	Trp	Xaa	Xaa	Xaa
	195					200						205			
Xaa	Gln	Lys	Xaa	Xaa	Ile	Val	Xaa	Xaa	Xaa	Xaa	Glu	Xaa	Lys	Lys	Met
210						215					220				
Thr	Phe	Asp	Xaa	Xaa	Xaa	Lys	Xaa	Xaa	Met	Gly	Xaa	Xaa	Pro	Xaa	Xaa
225					230					235					240
Glu	Xaa	Thr	Xaa	Xaa	Xaa	Xaa	Leu	Val	Xaa	Glu	Xaa	Glu	Xaa	Leu	Ile
				245					250					255	
Lys	Gly	Leu	Phe	Ser	Leu	Pro	Ile	Asn	Leu	Pro	Xaa	Thr	Ala	Tyr	Xaa
			260					265					270		
Lys	Ala	Leu	Xaa	Ala	Arg	Ala	Phe	Xaa	Xaa	Ala	Xaa	Leu	Glu	Xaa	Xaa
		275					280					285			
Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Ile	Xaa	Glu	Xaa	Arg	Xaa	Glu	Glu
290						295					300				
Glu	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa
305					310					315					320
Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Asp	Asp	Leu	Leu	Gly	Leu	Leu	Xaa	Ala	Xaa
				325					330					335	
Xaa	Xaa	Xaa	Xaa	Xaa	Glu	Asp	Glu	Xaa	Xaa	Xaa	Xaa	Leu	Ser	Asp	Xaa
			340					345					350		
Glu	Ile	Xaa	Asp	Xaa	Ile	Xaa	Xaa	Leu	Leu	Phe	Ala	Gly	His	Glu	Thr
	355						360					365			
Thr	Ser	Xaa	Xaa	Leu	Xaa	Xaa	Ala	Val	Lys	Phe	Leu	Xaa	Glu	His	Pro
	370					375					380				
Asp	Val	Xaa	Glu	Xaa	Leu	Arg	Glu	Glu	His	Xaa	Ala	Ile	Xaa	Arg	Ala
385					390					395					400
Lys	Lys	Xaa	Xaa	Xaa	Glu	Ser	Xaa	Leu	Thr	Xaa	Xaa	Asp	Tyr	Lys	Lys
				405					410					415	
Met	Xaa	Tyr	Thr	Xaa	Cys	Val	Ile	Asn	Glu	Thr	Leu	Arg	Leu	Ala	



Arg	Leu	Cys	Pro	Gly	Lys	Glu	Leu	Ala	Lys	Leu	Glu	Met	Xaa	Val	Phe
		515					520					525			
Leu	His	Arg	Leu	Val	Gln	Xaa	Phe	Trp	Glu	Leu	Ala	Xaa	Xaa	Xaa	Asp
	530					535					540				
Xaa	Xaa	Xaa	Lys	Leu	Val	Xaa	Phe	Pro	Thr	Xaa	Arg	Pro	Xaa	Asp	Asn
545					550					555					560
Leu	Pro	Ile	Lys	Val	Xaa	Xaa	Arg	Asp	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	
				565					570					575	